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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/10/036,492

DATE: 04/24/2002
 TIME: 14:28:08

Input Set : A:\217943US0XCONT.txt
 Output Set: N:\CRF3\04242002\J036492.raw

3 <110> APPLICANT: HEMERLY, ADRIANA
 4 FERREIRA, PAULO
 5 ROMBAUTS, STEPHANE
 7 <120> TITLE OF INVENTION: PLANT DNA REPLICATION MODULATING PROTEINS
 9 <130> FILE REFERENCE: 217943US0XCONT
 11 <140> CURRENT APPLICATION NUMBER: US 10/036,492
 12 <141> CURRENT FILING DATE: 2002-01-07
 14 <150> PRIOR APPLICATION NUMBER: EP 99202214.5
 15 <151> PRIOR FILING DATE: 1999-07-05
 17 <160> NUMBER OF SEQ ID NOS: 35
 19 <170> SOFTWARE: PatentIn version 3.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 889
 23 <212> TYPE: PRT
 24 <213> ORGANISM: Arabidopsis thaliana
 26 <400> SEQUENCE: 1
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 29 1 5 10 15
 32 Arg Glu Leu Ile Pro Leu Ser Pro Thr Asn Ser Asp Gly Asn Asp Asp
 33 20 25 30
 36 Leu Asn Tyr His Leu His Ala Phe Glu Leu Ser Arg Leu Leu Leu Ser
 37 35 40 45
 40 Ser Gly His Pro Glu Ser Val Ile Asp Leu Ser Ser Lys Cys Thr Tyr
 41 50 55 60
 44 Phe Gln Gly Ser Pro Asn Leu Val Lys Tyr Leu Cys Ser Ile Pro Asn
 45 65 70 75 80
 48 Ser Pro Ile Ser Leu Ala Glu Asp Gly Phe Thr Val Thr Leu Ser Pro
 49 85 90 95
 52 Glu Ser Pro Ser Ala Pro Ala Ser Phe Ala Cys Ser Leu Asp Leu Gln
 53 100 105 110
 56 Glu Asn Val Val Leu Glu Gln Phe Met Asp Pro Arg Ser Leu Thr Leu
 57 115 120 125
 60 Lys His Ser Arg Glu Asn Ala Glu Gln Glu Leu Glu Leu Met Pro
 61 130 135 140
 64 Leu Pro Lys Arg Ser Arg Asn Asp Gly Asn Asp Val Asn Tyr Ser Val
 65 145 150 155 160
 68 Ile Asp Ser Arg Pro Asn Asp Ile Arg Thr Val Ala Cys Gly Thr Met
 69 165 170 175
 72 Leu Gly Thr Ile Leu Ala Leu Glu Ser Gln Ala Ser Val Phe Asn Leu
 73 180 185 190
 76 Ser Ala Ser Asn Arg Gly Ile Glu Ala Phe Val Gln Asp His Gln Pro
 77 195 200 205
 80 Gly Pro Gln Thr Ser Asn Ala Ser Val Asp Val Asn Pro Thr His Arg

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81	210	215	220
84	Leu Glu Glu Ser Lys Asn Asp Leu Pro Ser Pro Gln Glu Asp Gly Tyr		
85	225	230	235
88	Tyr Glu Arg Pro Glu Ile Gly Asp Phe Gln Ile Ala Asp Asn Gln Ile		240
89	245	250	255
92	Leu Ile Glu Glu Gly Asp Asp Lys Asn Lys Lys Asp Leu Phe Pro Lys		
93	260	265	270
96	Gly Glu Ile Gln Thr Asp Ser Val Gln Ser Asp Pro Val Ala Ser Leu		
97	275	280	285
100	Met Pro Thr Glu Asn Glu Leu Glu Pro Val Gln Ile Val Asp Asp Thr		
101	290	295	300
104	Glu Asp Leu Leu Val Asp Asp His Thr Val Asp Ile Val Ser Thr Pro		
105	305	310	315
108	Asp Arg Glu Leu Pro Leu Lys Pro Ser Ala Thr Glu Ala Asn Gln Asp		320
109	325	330	335
112	Lys Ser Leu Val Gln Lys Thr Leu Asp Gln Cys Lys Leu Pro Gly Asn		
113	340	345	350
116	Ser Lys Thr Tyr Ser Cys Ser Pro Glu Ile Lys His Thr Arg Lys Ser		
117	355	360	365
120	Lys Val Ile Gln Lys Arg Lys Gln Asn Phe Asn Thr Val Arg Leu Lys		
121	370	375	380
124	Asp Gln Lys Asp Gln Ala Lys His Asn Thr Ile Pro Asp Phe Asp Ser		
125	385	390	395
128	Tyr Thr Ile Val Glu Glu Gly Ser Gly Gly Tyr Gly Ile Val Tyr		
129	405	410	415
132	Lys Ala Thr Arg Lys Thr Asp Gly Thr Glu Phe Ala Ile Lys Cys Pro		
133	420	425	430
136	His Val Gly Ala Gln Lys Tyr Tyr Val Asn Asn Glu Ile Arg Met Leu		
137	435	440	445
140	Glu Arg Phe Gly Gly Lys Asn Cys Ile Ile Lys His Glu Gly Cys Leu		
141	450	455	460
144	Lys Asn Gly Asp Ser Asp Cys Ile Ile Leu Glu His Leu Glu His Asp		
145	465	470	475
148	Arg Pro Asp Ser Leu Lys Arg Glu Ile Asp Val Tyr Gln Leu Gln Trp		480
149	485	490	495
152	Tyr Gly Tyr Cys Met Phe Lys Ala Leu Ser Ser Leu His Lys Gln Gly		
153	500	505	510
156	Val Val His Arg Asp Val Lys Pro Gly Asn Phe Leu Phe Ser Arg Lys		
157	515	520	525
160	Thr Asn Lys Gly Tyr Leu Ile Asp Phe Asn Leu Ala Met Asp Leu His		
161	530	535	540
164	Gln Lys Tyr Arg Arg Ala Asp Lys Ser Lys Ala Ala Ser Gly Leu Pro		
165	545	550	555
168	Thr Ala Ser Lys Lys His His Thr Leu Val Lys Ser Leu Asp Ala Val		560
169	565	570	575
172	Asn Arg Gly Thr Asn Lys Pro Ser Gln Lys Thr Leu Ala Pro Asn Ser		
173	580	585	590
176	Ile Lys Lys Ala Ala Gly Lys Thr Arg Ala Arg Asn Asp Met Thr Arg		
177	595	600	605

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180 Trp Glu Arg Leu Asn Ser Gln Gly Ala Glu Gly Ser Gly Leu Thr Ser
181 610 615 620
184 Ala Lys Asp Val Thr Ser Thr Arg Asn Asn Pro Ser Gly Glu Lys Arg
185 625 630 635 640
188 Arg Glu Pro Leu Pro Cys His Gly Arg Lys Ala Leu Leu Asp Phe Leu
189 645 650 655
192 Gln Glu Thr Met Ser Val Pro Ile Pro Asn His Glu Val Ser Ser Lys
193 660 665 670
196 Ala Pro Thr Ser Met Arg Lys Arg Val Ala Ala Leu Pro Gly Lys Ala
197 675 680 685
200 Glu Lys Glu Leu Leu Tyr Leu Thr Pro Met Pro Leu Cys Ser Asn Gly
201 690 695 700
204 Arg Pro Glu Ala Gly Asp Val Ile Glu Lys Lys Asp Gly Pro Cys Ser
205 705 710 715 720
208 Gly Thr Lys Gly Phe Arg Ala Pro Glu Val Cys Phe Arg Ser Leu His
209 725 730 735
212 Gln Gly Pro Lys Ile Asp Val Trp Ser Ala Gly Val Thr Leu Leu Tyr
213 740 745 750
216 Leu Ile Met Gly Arg Thr Pro Phe Thr Gly Asp Pro Glu Gln Asn Ile
217 755 760 765
220 Lys Asp Ile Ala Gln Leu Arg Gly Ser Glu Glu Leu Trp Glu Val Ala
221 770 775 780
224 Lys Leu His Asn Arg Glu Ser Ser Phe Pro Lys Glu Leu Tyr Glu Ser
225 785 790 795 800
228 Arg Tyr Leu Lys Gly Met Glu Leu Arg Lys Trp Cys Glu Leu Asn Thr
229 805 810 815
232 Lys Arg Arg Glu Phe Leu Asp Val Ile Pro Leu Ser Leu Leu Asp Leu
233 820 825 830
236 Val Asp Lys Cys Leu Thr Val Asn Pro Arg Arg Arg Ile Ser Ala Glu
237 835 840 845
240 Asp Ala Leu Lys His Asp Phe Phe His Pro Val His Glu Thr Leu Arg
241 850 855 860
244 Asn Gln Met Leu Leu Lys Gln Gln Pro Thr Val Val Ala Asp Ala Val
245 865 870 875 880
248 Ser Gln Thr Leu Asn Tyr Leu Gln Leu
249 885
252 <210> SEQ ID NO: 2
253 <211> LENGTH: 20
254 <212> TYPE: PRT
255 <213> ORGANISM: Artificial Sequence
257 <220> FEATURE:
258 <223> OTHER INFORMATION: Synthetic Peptide
260 <400> SEQUENCE: 2
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263 1 5 10 15
266 Phe Ala Ile Lys
267 20
270 <210> SEQ ID NO: 3
271 <211> LENGTH: 20

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Input Set : A:\217943US0XCONT.txt
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272 <212> TYPE: PRT
 273 <213> ORGANISM: Artificial Sequence
 275 <220> FEATURE:
 276 <223> OTHER INFORMATION: Synthetic Peptide
 278 <400> SEQUENCE: 3
 280 Asp Val Ile Glu Lys Lys Asp Gly Pro Cys Ser Gly Thr Lys Gly Phe
 281 1 5 10 15
 284 Arg Ala Pro Glu
 285 20
 288 <210> SEQ ID NO: 4
 289 <211> LENGTH: 29
 290 <212> TYPE: PRT
 291 <213> ORGANISM: Artificial Sequence
 293 <220> FEATURE:
 294 <223> OTHER INFORMATION: Synthetic Peptide
 296 <400> SEQUENCE: 4
 298 Asn Ile Lys Asp Ile Ala Gln Leu Arg Gly Ser Glu Glu Leu Trp Glu
 299 1 5 10 15
 302 Val Ala Lys Leu His Asn Arg Glu Ser Ser Phe Pro Lys
 303 20 25
 306 <210> SEQ ID NO: 5
 307 <211> LENGTH: 728
 308 <212> TYPE: PRT
 309 <213> ORGANISM: Arabidopsis thaliana
 311 <400> SEQUENCE: 5
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 314 1 5 10 15
 317 Phe Met Phe Thr Asn Ala Ile Phe Leu Cys Glu Leu Leu Leu Ala Gln
 318 20 25 30
 321 Phe Pro Ser Glu Val Asn Leu Gln Leu Leu Ala Arg Cys Tyr Leu Ser
 322 35 40 45
 325 Asn Ser Gln Ala Tyr Ser Ala Tyr Tyr Ile Leu Lys Gly Ser Lys Thr
 326 50 55 60
 329 Pro Gln Ser Arg Tyr Leu Phe Ala Phe Ser Cys Phe Lys Leu Asp Leu
 330 65 70 75 80
 333 Leu Gly Glu Ala Glu Ala Ala Leu Leu Pro Cys Glu Asp Tyr Ala Glu
 334 85 90 95
 337 Glu Val Pro Gly Gly Ala Ala Gly His Tyr Leu Leu Gly Leu Ile Tyr
 338 100 105 110
 341 Arg Tyr Ser Gly Arg Lys Asn Cys Ser Ile Gln Gln Phe Arg Met Ala
 342 115 120 125
 345 Leu Ser Phe Asp Pro Leu Cys Trp Glu Ala Tyr Gly Glu Leu Cys Ser
 346 130 135 140
 349 Leu Gly Ala Ala Glu Glu Ala Ser Thr Val Phe Gly Asn Val Ala Ser
 350 145 150 155 160
 353 Gln Arg Leu Gln Lys Thr Cys Val Glu Gln Arg Ile Ser Phe Ser Glu
 354 165 170 175
 357 Gly Ala Thr Ile Asp Gln Ile Thr Asp Ser Asp Lys Ala Leu Lys Asp
 358 180 185 190

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Input Set : A:\217943US0XCONT.txt
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361 Thr Gly Leu Ser Gln Thr Glu His Ile Pro Gly Glu Asn Gln Gln Asp
 362 195 200 205
 365 Leu Lys Ile Met Gln Gln Pro Gly Asp Ile Pro Pro Asn Thr Asp Arg
 366 210 215 220
 369 Gln Leu Ser Thr Asn Gly Trp Asp Leu Asn Thr Pro Ser Pro Val Leu
 370 225 230 235 240
 373 Leu Gln Val Met Asp Ala Leu Pro Pro Leu Leu Leu Lys Asn Met Arg
 374 245 250 255
 377 Arg Pro Ala Val Glu Gly Ser Leu Met Ser Val His Gly Val Arg Val
 378 260 265 270
 381 Arg Arg Arg Asn Phe Phe Ser Glu Glu Leu Ser Ala Glu Ala Gln Glu
 382 275 280 285
 385 Glu Ser Gly Arg Arg Arg Ser Ala Arg Ile Ala Ala Arg Lys Lys Asn
 386 290 295 300
 389 Pro Met Ser Gln Ser Phe Gly Lys Asp Ser His Trp Leu His Leu Ser
 390 305 310 315 320
 393 Pro Ser Glu Ser Asn Tyr Ala Pro Ser Leu Ser Ser Met Ile Gly Lys
 394 325 330 335
 397 Cys Arg Ile Gln Ser Ser Lys Glu Val Ile Pro Asp Thr Val Thr Leu
 398 340 345 350
 401 Asn Asp Pro Ala Thr Thr Ser Gly Gln Ser Val Ser Asp Ile Gly Ser
 402 355 360 365
 405 Ser Val Asp Asp Glu Glu Lys Ser Asn Pro Ser Glu Ser Ser Pro Asp
 406 370 375 380
 409 Arg Phe Ser Leu Ile Ser Gly Ile Ser Glu Val Leu Ser Leu Leu Lys
 410 385 390 395 400
 413 Ile Leu Gly Asp Gly His Arg His Leu His Met Tyr Lys Cys Gln Glu
 414 405 410 415
 417 Ala Leu Leu Ala Tyr Gln Lys Leu Ser Gln Lys Gln Tyr Asn Thr His
 418 420 425 430
 421 Trp Val Leu Met Gln Val Gly Lys Ala Tyr Phe Glu Leu Gln Asp Tyr
 422 435 440 445
 425 Phe Asn Ala Asp Ser Ser Phe Thr Leu Ala His Gln Lys Tyr Pro Tyr
 426 450 455 460
 429 Ala Leu Glu Gly Met Asp Thr Tyr Ser Thr Val Leu Tyr His Leu Lys
 430 465 470 475 480
 433 Glu Glu Met Arg Leu Gly Tyr Leu Ala Gln Glu Leu Ile Ser Val Asp
 434 485 490 495
 437 Arg Leu Ser Pro Glu Ser Trp Cys Ala Val Gly Asn Cys Tyr Ser Leu
 438 500 505 510
 441 Arg Lys Asp His Asp Thr Ala Leu Lys Met Phe Gln Arg Ala Ile Gln
 442 515 520 525
 445 Leu Asn Glu Arg Phe Thr Tyr Ala His Thr Leu Cys Gly His Glu Phe
 446 530 535 540
 449 Ala Ala Leu Glu Glu Phe Glu Asp Ala Glu Arg Cys Tyr Arg Lys Ala
 450 545 550 555 560
 453 Leu Gly Ile Asp Thr Arg His Tyr Asn Ala Trp Tyr Gly Leu Gly Met
 454 565 570 575
 457 Thr Tyr Leu Arg Gln Glu Lys Phe Glu Phe Ala Gln His Gln Phe Gln

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/24/2002
PATENT APPLICATION: US/10/036,492 TIME: 14:28:10

Input Set : A:\217943US0XCONT.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:16; Xaa Pos. 1,2,3,5,6,9,10,12,13,14,15,16,17,18,19,21,22,23,25,26,28
Seq#:16; Xaa Pos. 29,30,31,33,34
Seq#:18; N Pos. 9,15,18
Seq#:20; N Pos. 6,9,18
Seq#:21; N Pos. 3,6,15
Seq#:32; N Pos. 6,12
Seq#:33; N Pos. 6,15
Seq#:34; N Pos. 15
Seq#:35; N Pos. 9